
Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: [year=2010; month=5; day=3; hr=10; min=10; sec=43; ms=829;]

Validated By CRFValidator v 1.0.3

Application No: 10590870 Version No: 1.0

Input Set:

Output Set:

Started: 2010-04-23 14:53:05.942 **Finished:** 2010-04-23 14:53:09.612

Elapsed: 0 hr(s) 0 min(s) 3 sec(s) 670 ms

Total Warnings: 51
Total Errors: 2
No. of SeqIDs Defined: 51

Actual SeqID Count: 51

Error code		Error Description
W	213	Artificial or Unknown found in <213> in SEQ ID (1)
E	355	Empty lines found between the amino acid numbering and the
E	321	No. of Bases conflict, this line has no nucleotides SEQID (1)
W	213	Artificial or Unknown found in <213> in SEQ ID (2)
W	213	Artificial or Unknown found in <213> in SEQ ID (3)
W	213	Artificial or Unknown found in <213> in SEQ ID (4)
W	213	Artificial or Unknown found in <213> in SEQ ID (5)
W	213	Artificial or Unknown found in <213> in SEQ ID (6)
W	213	Artificial or Unknown found in <213> in SEQ ID (7)
W	213	Artificial or Unknown found in <213> in SEQ ID (8)
W	213	Artificial or Unknown found in <213> in SEQ ID (9)
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W	213	Artificial or Unknown found in <213> in SEQ ID (17)
W	213	Artificial or Unknown found in <213> in SEQ ID (18)

Input Set:

Output Set:

Started: 2010-04-23 14:53:05.942 **Finished:** 2010-04-23 14:53:09.612

Elapsed: 0 hr(s) 0 min(s) 3 sec(s) 670 ms

Total Warnings: 51
Total Errors: 2

No. of SeqIDs Defined: 51

Actual SeqID Count: 51

Error code		Error Description
W	213	Artificial or Unknown found in <213> in SEQ ID (19)
W	213	Artificial or Unknown found in <213> in SEQ ID (20) This error has occured more than 20 times, will not be displayed

SEQUENCE LISTING

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     Harding, Fiona A.
      Rashid, M. Harunur
      Schellenberger, Volker
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Ser Val Thr Val Ser Ser Thr Pro Val Ser Glu Lys Gln Leu Ala Glu 260 265 Val Val Ala Asn Thr Ile Thr Pro Leu Met Lys Ala Gln Ser Val Pro 280 Gly Met Ala Val Ala Val Ile Tyr Gln Gly Lys Pro His Tyr Tyr Thr 295 300 Phe Gly Lys Ala Asp Ile Ala Ala Asn Lys Pro Val Thr Pro Gln Thr 310 315 320 Leu Phe Glu Leu Gly Ser Ile Ser Lys Thr Phe Thr Gly Val Leu Gly 325 330 Gly Asp Ala Ile Ala Arg Gly Glu Ile Ser Leu Asp Asp Ala Val Thr 340 345 350 Arg Tyr Trp Pro Gln Leu Thr Gly Lys Gln Trp Gln Gly Ile Arg Met 360 Leu Asp Leu Ala Thr Tyr Thr Ala Gly Gly Leu Pro Leu Gln Val Pro 375 380 Asp Glu Val Thr Asp Asn Ala Ser Leu Leu Arg Phe Tyr Gln Asn Trp 390 385 395 400 Gln Pro Gln Trp Lys Pro Gly Thr Thr Arg Leu Tyr Ala Asn Ala Ser 405 410 Ile Gly Leu Phe Gly Ala Leu Ala Val Lys Pro Ser Gly Met Pro Tyr 420 425 Glu Gln Ala Met Thr Thr Arg Val Leu Lys Pro Leu Lys Leu Asp His 440 Thr Trp Ile Asn Val Pro Lys Ala Glu Glu Ala His Tyr Ala Trp Gly 455 460 Tyr Arg Asp Gly Lys Ala Val Arg Val Ser Pro Gly Met Leu Asp Ala 470 475 480 Gln Ala Tyr Gly Val Lys Thr Asn Val Gln Asp Met Ala Asn Trp Val 485 490 Met Ala Asn Met Ala Pro Glu Asn Val Ala Asp Ala Ser Leu Lys Gln 505 500 Gly Ile Ala Leu Ala Gln Ser Arg Tyr Trp Arg Ile Gly Ser Met Tyr 515 520 525 Gln Gly Leu Gly Trp Glu Met Leu Asn Trp Pro Val Glu Ala Asn Thr 535 Val Val Glu Thr Ser Phe Gly Asn Val Ala Leu Ala Pro Leu Pro Val 550 555 560 Ala Glu Val Asn Pro Pro Ala Pro Pro Val Lys Ala Ser Trp Val His 565 570 575 Lys Thr Gly Ser Thr Gly Gly Phe Gly Ser Tyr Val Ala Phe Ile Pro 585 Glu Lys Gln Ile Gly Ile Val Met Leu Ala Asn Thr Ser Tyr Pro Asn 600 Pro Ala Arg Val Glu Ala Ala Tyr His Ile Leu Glu Ala Leu Gln 615 <210> 3 <211> 786 <212> DNA <213> Artificial Sequence

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1869

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420

480 540

600

660 720

780 786 290 295 300

Phe Gly Lys Ala Asp Ile Ala Ala Asn Lys Pro Val Thr Pro Gln Thr 305 310 315 320

Leu Phe Glu Leu Gly Ser Ile Ser Lys Thr Phe Thr Gly Val L